

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Application of: RADIN et al.

Appln. No.: 08/713,928

Group Art Unit: 1646

Filed: September 13, 1996

Examiner: E. KEMMERER

For: PRODUCTION OF LYSOSOMAL
ENZYMES IN PLANT-BASED
EXPRESSION SYSTEMS

Atty. Dkt. No.: 7956-011

DECLARATION UNDER 37 C.F.R. § 1.132

Assistant Commissioner for Patents
Washington, D.C. 20231

Sir:

I, KAREN K. OISHI, declare and state that:

1. I am a co-inventor of the above-identified application.

2. I have carefully reviewed autoradiograms containing sequence data of the MeGA promoter and compared that data with the MeGA promoter sequence shown in Figure 11 originally filed with the above-identified application.

3. My review indicates that several nucleotide residues were inadvertently omitted from or incorrectly recorded in the sequence shown in the originally filed Figure 11.

4. The correct sequence of the MeGA promoter is shown in the hand-corrected Figure 11 attached hereto as Exhibit A.

5. The correct MeGA promoter sequence was based on data obtained from a cloned MeGA promoter-containing DNA that was also used in constructing the MeGA:IDUA expression construct contained in plasmid pCT22, which was deposited with American Type Culture Collection on August 30, 1996 and assigned accession number ATCC 97701.

I hereby declare further that all statements made herein of my own knowledge are true and that all statements made on information and belief are believed to be true; and further that these statements were made with the knowledge that willful false statements and the like so made are punishable by fine or imprisonment, or both, under § 1001 of Title 18 of the United States Code and that such willful false statements may jeopardize the validity of the application or any patent issuing thereon.

Date: 8/13/98

Karen K. Oishi
KAREN K. OISHI

CAATACGATA TTACCGAATA TTATACTAAA TCAAAATTTA ATTTATCATA TCAATTATTA 60
 AACTGATATT TCAAAATTTA ATATTTAATA TCTACTTTCA ACTATTATTA CCTAATTATC 120
 AAATGCAAAA TGTATGAGTT ATTTCATAAT AGCCCAGTTC GTATCCAAAT ATTTTACACT 180
 TGACCAGTCA ACTTGACTAT ATAAAACTTT ACTTCAAAAA ATTAAAAAAA AAAGAAAGTA 240
 TATTATTGTA AAAGATAATA CTCCATTCAA AATATAAAAT GAAAAAAGTC CAGCGCGGCA 300
 ACCGGGTTCC TATAAATACA TTTCCTACAT CTTCTCTTCT CCTCACATCC CATCACTCTT 360
 CTTTAAACAA TTATACTTGT CAATCATCAA TCCCACAAAC AACACTTTTT CTCTCCTCTT 420
 TTTCCTCACC GCGGCAGAC TTACCGGTGA AAGTAAGCAG⁴⁶³ CTC
 TCTAGA

FIG. 11